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Reviewer: Keisha Douglas

Timestamp: [year=2008; month=9; day=10; hr=15; min=10; sec=15; ms=998;]

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Application No: 10598682

Version No: 1.0

Input Set:**Output Set:****Started:** 2008-08-11 11:17:24.606**Finished:** 2008-08-11 11:17:26.881**Elapsed:** 0 hr(s) 0 min(s) 2 sec(s) 275 ms**Total Warnings:** 22**Total Errors:** 0**No. of SeqIDs Defined:** 57**Actual SeqID Count:** 57

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Input Set:

Output Set:

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Total Errors: 0
No. of SeqIDs Defined: 57
Actual SeqID Count: 57

Error code

Error Description

This error has occurred more than 20 times, will not be displayed

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CELLS

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<140> 10598682

<141> 2008-08-11

<150> PCT/US05/007704

<151> 2005-03-09

<150> 60/552,318

<151> 2004-03-10

<160> 57

<170> PatentIn Ver. 3.3

<210> 1

<211> 116

<212> PRT

<213> Homo sapiens

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			20					25						30	

Ser	Gly	Tyr	His	Ala	Asn	Tyr	Cys	Glu	Gly	Glu	Cys	Pro	Ser	His	Ile
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Ala	Gly	Thr	Ser	Gly	Ser	Ser	Leu	Ser	Phe	His	Ser	Thr	Val	Ile	Asn
			50				55				60				

His	Tyr	Arg	Met	Arg	Gly	His	Ser	Pro	Phe	Ala	Asn	Leu	Lys	Ser	Cys
65					70				75					80	

Cys	Val	Pro	Thr	Lys	Leu	Arg	Pro	Met	Ser	Met	Leu	Tyr	Tyr	Asp	Asp
				85				90						95	

Gly	Gln	Asn	Ile	Ile	Lys	Lys	Asp	Ile	Gln	Asn	Met	Ile	Val	Glu	Glu
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<213> Homo sapiens

<400> 2

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gaggggtgagt gcccagagcca tatagcaggc acgtccgggt cctcactgtc cttccactca 180
acagtcatca accactaccg catgcggggc catagcccct ttgccaacct caaatcgtgc 240
tgtgtgccca ccaagctgag acccatgtcc atgttgtact atgatgatgg tcaaaacatc 300
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<212> PRT

<213> Homo sapiens

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Ile Val Arg Ser Ser Pro Thr Pro Gly Ser Glu Gly His Ser Ala Ala
      20              25              30
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Pro Asp Cys Pro Ser Cys Ala Leu Ala Ala Leu Pro Lys Asp Val Pro
      35              40              45
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```
Asn Ser Gln Pro Glu Met Val Glu Ala Val Lys Lys His Ile Leu Asn
      50              55              60
```

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Met Leu His Leu Lys Lys Arg Pro Asp Val Thr Gln Pro Val Pro Lys
      65              70              75              80
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Ala Ala Leu Leu Asn Ala Ile Arg Lys Leu His Val Gly Lys Val Gly
      85              90              95
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Glu Asn Gly Tyr Val Glu Ile Glu Asp Asp Ile Gly Arg Arg Ala Glu
      100             105             110
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Met Asn Glu Leu Met Glu Gln Thr Ser Glu Ile Ile Thr Phe Ala Glu
      115             120             125
```

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Ser Gly Thr Ala Arg Lys Thr Leu His Phe Glu Ile Ser Lys Glu Gly
      130             135             140
```

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Ser Asp Leu Ser Val Val Glu Arg Ala Glu Val Trp Leu Phe Leu Lys
      145             150             155             160
```

```
Val Pro Lys Ala Asn Arg Thr Arg Thr Lys Val Thr Ile Arg Leu Phe
      165             170             175
```

```
Gln Gln Gln Lys His Pro Gln Gly Ser Leu Asp Thr Gly Glu Glu Ala
      180             185             190
```

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Glu Glu Val Gly Leu Lys Gly Glu Arg Ser Glu Leu Leu Leu Ser Glu
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Lys Val Val Asp Ala Arg Lys Ser Thr Trp His Val Phe Pro Val Ser
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Ser Ser Ile Gln Arg Leu Leu Asp Gln Gly Lys Ser Ser Leu Asp Val
225 230 235 240

Arg Ile Ala Cys Glu Gln Cys Gln Glu Ser Gly Ala Ser Leu Val Leu
245 250 255

Leu Gly Lys Lys Lys Lys Lys Glu Glu Glu Gly Glu Gly Lys Lys Lys
260 265 270

Gly Gly Gly Glu Gly Gly Ala Gly Ala Asp Glu Glu Lys Glu Gln Ser
275 280 285

His Arg Pro Phe Leu Met Leu Gln Ala Arg Gln Ser Glu Asp His Pro
290 295 300

His Arg Arg Arg Arg Arg Gly Leu Glu Cys Asp Gly Lys Val Asn Ile
305 310 315 320

Cys Cys Lys Lys Gln Phe Phe Val Ser Phe Lys Asp Ile Gly Trp Asn
325 330 335

Asp Trp Ile Ile Ala Pro Ser Gly Tyr His Ala Asn Tyr Cys Glu Gly
340 345 350

Glu Cys Pro Ser His Ile Ala Gly Thr Ser Gly Ser Ser Leu Ser Phe
355 360 365

His Ser Thr Val Ile Asn His Tyr Arg Met Arg Gly His Ser Pro Phe
370 375 380

Ala Asn Leu Lys Ser Cys Cys Val Pro Thr Lys Leu Arg Pro Met Ser
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Asn Met Ile Val Glu Glu Cys Gly Cys Ser
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<210> 4

<211> 130

<212> PRT

<213> Homo sapiens

<400> 4

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20 25 30

Ser Phe Lys Asp Ile Gly Trp Asn Asp Trp Ile Ile Ala Pro Ser Gly
35 40 45

Tyr His Ala Asn Tyr Cys Glu Gly Glu Cys Pro Ser His Ile Ala Gly

50

55

60

Thr Ser Gly Ser Ser Leu Ser Phe His Ser Thr Val Ile Asn His Tyr
65 70 75 80

Arg Met Arg Gly His Ser Pro Phe Ala Asn Leu Lys Ser Cys Cys Val
85 90 95

Pro Thr Lys Leu Arg Pro Met Ser Met Leu Tyr Tyr Asp Asp Gly Gln
100 105 110

Asn Ile Ile Lys Lys Asp Ile Gln Asn Met Ile Val Glu Glu Cys Gly
115 120 125

Cys Ser
130

<210> 5

<211> 115

<212> PRT

<213> Homo sapiens

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Phe Val Ser Phe Lys Asp Ile Gly Trp Asn Asp Trp Ile Ile Ala Pro
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Ser Gly Tyr His Ala Asn Tyr Cys Glu Gly Glu Cys Pro Ser His Ile
35 40 45

Ala Gly Thr Ser Gly Ser Ser Leu Ser Phe His Ser Thr Val Ile Asn
50 55 60

His Tyr Ala Cys Gly His Ser Pro Phe Ala Asn Leu Lys Ser Cys Cys
65 70 75 80

Val Pro Thr Lys Leu Arg Pro Met Ser Met Leu Tyr Tyr Asp Asp Gly
85 90 95

Gln Asn Ile Ile Lys Lys Asp Ile Gln Asn Met Ile Val Glu Glu Cys
100 105 110

Gly Cys Ser
115

<210> 6

<211> 426

<212> PRT

<213> Homo sapiens

<400> 6

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			20					25					30				
Pro	Asp	Cys	Pro	Ser	Cys	Ala	Leu	Ala	Ala	Leu	Pro	Lys	Asp	Val	Pro		
		35					40					45					
Asn	Ser	Gln	Pro	Glu	Met	Val	Glu	Ala	Val	Lys	Lys	His	Ile	Leu	Asn		
		50				55						60					
Met	Leu	His	Leu	Lys	Lys	Arg	Pro	Asp	Val	Thr	Gln	Pro	Val	Pro	Lys		
		65			70					75					80		
Ala	Ala	Leu	Leu	Asn	Ala	Ile	Arg	Lys	Leu	His	Val	Gly	Lys	Val	Gly		
				85					90					95			
Glu	Asn	Gly	Tyr	Val	Glu	Ile	Glu	Asp	Asp	Ile	Gly	Arg	Arg	Ala	Glu		
			100					105					110				
Met	Asn	Glu	Leu	Met	Glu	Gln	Thr	Ser	Glu	Ile	Ile	Thr	Phe	Ala	Glu		
		115					120					125					
Ser	Gly	Thr	Ala	Arg	Lys	Thr	Leu	His	Phe	Glu	Ile	Ser	Lys	Glu	Gly		
		130				135						140					
Ser	Asp	Leu	Ser	Val	Val	Glu	Arg	Ala	Glu	Val	Trp	Leu	Phe	Leu	Lys		
		145			150					155					160		
Val	Pro	Lys	Ala	Asn	Arg	Thr	Arg	Thr	Lys	Val	Thr	Ile	Arg	Leu	Phe		
				165					170					175			
Gln	Gln	Gln	Lys	His	Pro	Gln	Gly	Ser	Leu	Asp	Thr	Gly	Glu	Glu	Ala		
			180					185					190				
Glu	Glu	Val	Gly	Leu	Lys	Gly	Glu	Arg	Ser	Glu	Leu	Leu	Leu	Ser	Glu		
		195					200					205					
Lys	Val	Val	Asp	Ala	Arg	Lys	Ser	Thr	Trp	His	Val	Phe	Pro	Val	Ser		
		210				215					220						
Ser	Ser	Ile	Gln	Arg	Leu	Leu	Asp	Gln	Gly	Lys	Ser	Ser	Leu	Asp	Val		
		225			230					235				240			
Arg	Ile	Ala	Cys	Glu	Gln	Cys	Gln	Glu	Ser	Gly	Ala	Ser	Leu	Val	Leu		
			245					250					255				
Leu	Gly	Lys	Lys	Lys	Lys	Lys	Glu	Glu	Glu	Gly	Glu	Gly	Lys	Lys	Lys		
			260					265					270				
Gly	Gly	Gly	Glu	Gly	Gly	Ala	Gly	Ala	Asp	Glu	Glu	Lys	Glu	Gln	Ser		
		275					280					285					
His	Arg	Pro	Phe	Leu	Met	Leu	Gln	Ala	Arg	Gln	Ser	Glu	Asp	His	Pro		
		290				295					300						
His	Arg	Arg	Arg	Arg	Arg	Gly	Leu	Glu	Cys	Asp	Gly	Lys	Val	Asn	Ile		
		305			310					315				320			

Cys Cys Lys Lys Gln Phe Phe Val Ser Phe Lys Asp Ile Gly Trp Asn
 325 330 335

Asp Trp Ile Ile Ala Pro Ser Gly Tyr His Ala Asn Tyr Cys Glu Gly
 340 345 350

Glu Cys Pro Ser His Ile Ala Gly Thr Ser Gly Ser Ser Leu Ser Phe
 355 360 365

His Ser Thr Val Ile Asn His Tyr Arg Met Arg Gly His Ser Pro Phe
 370 375 380

Ala Asn Leu Lys Ser Cys Cys Val Pro Thr Lys Leu Arg Pro Met Ser
 385 390 395 400

Met Leu Tyr Tyr Asp Asp Gly Gln Asn Ile Ile Lys Lys Asp Ile Gln
 405 410 415

Asn Met Ile Val Glu Glu Cys Gly Cys Ser
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 <212> PRT
 <213> Mus musculus

<400> 7
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 20 25 30

Pro Asp Cys Pro Ser Cys Ala Leu Ala Thr Leu Pro Lys Asp Gly Pro
 35 40 45

Asn Ser Gln Pro Glu Met Val Glu Ala Val Lys Lys His Ile Leu Asn
 50 55 60

Met Leu His Leu Lys Lys Arg Pro Asp Val Thr Gln Pro Val Pro Lys
 65 70 75 80

Ala Ala Leu Leu Asn Ala Ile Arg Lys Leu His Val Gly Lys Val Gly
 85 90 95

Glu Asn Gly Tyr Val Glu Ile Glu Asp Asp Ile Gly Arg Arg Ala Glu
 100 105 110

Met Asn Glu Leu Met Glu Gln Thr Ser Glu Ile Ile Thr Phe Ala Glu
 115 120 125

Ser Gly Thr Ala Arg Lys Thr Leu His Phe Glu Ile Ser Lys Glu Gly
 130 135 140

Ser Asp Leu Ser Val Val Glu Arg Ala Glu Val Trp Leu Phe Leu Lys

145		150		155		160									
Val	Pro	Lys	Ala	Asn	Arg	Thr	Arg	Thr	Lys	Val	Thr	Ile	Arg	Leu	Phe
				165					170					175	
Gln	Gln	Gln	Lys	His	Pro	Gln	Gly	Ser	Leu	Asp	Thr	Gly	Asp	Glu	Ala
			180					185					190		
Glu	Glu	Met	Gly	Leu	Lys	Gly	Glu	Arg	Ser	Glu	Leu	Leu	Leu	Ser	Glu
		195					200					205			
Lys	Val	Val	Asp	Ala	Arg	Lys	Ser	Thr	Trp	His	Ile	Phe	Pro	Val	Ser
	210					215					220				
Ser	Ser	Ile	Gln	Arg	Leu	Leu	Asp	Gln	Gly	Lys	Ser	Ser	Leu	Asp	Val
225					230					235					240
Arg	Ile	Ala	Cys	Glu	Gln	Cys	Gln	Glu	Ser	Gly	Ala	Ser	Leu	Val	Leu
			245						250					255	
Leu	Gly	Lys	Lys	Lys	Lys	Lys	Glu	Val	Asp	Gly	Asp	Gly	Lys	Lys	Lys
		260						265					270		
Asp	Gly	Ser	Asp	Gly	Gly	Leu	Glu	Glu	Glu	Lys	Glu	Gln	Ser	His	Arg
	275						280					285			
Pro	Phe	Leu	Met	Leu	Gln	Ala	Arg	Gln	Ser	Glu	Asp	His	Pro	His	Arg
	290					295						300			
Arg	Arg	Arg	Arg	Gly	Leu	Glu	Cys	Asp	Gly	Lys	Val	Asn	Ile	Cys	Cys
305					310					315					320
Lys	Lys	Gln	Phe	Phe	Val	Ser	Phe	Lys	Asp	Ile	Gly	Trp	Asn	Asp	Trp
			325						330					335	
Ile	Ile	Ala	Pro	Ser	Gly	Tyr	His	Ala	Asn	Tyr	Cys	Glu	Gly	Glu	Cys
		340						345					350		
Pro	Ser	His	Ile	Ala	Gly	Thr	Ser	Gly	Ser	Ser	Leu	Ser	Phe	His	Ser
		355					360					365			
Thr	Val	Ile	Asn	His	Tyr	Arg	Met	Arg	Gly	His	Ser	Pro	Phe	Ala	Asn
	370					375					380				
Leu	Lys	Ser	Cys	Cys	Val	Pro	Thr	Lys	Leu	Arg	Pro	Met	Ser	Met	Leu
385					390					395					400
Tyr	Tyr	Asp	Asp	Gly	Gln	Asn	Ile	Ile	Lys	Lys	Asp	Ile	Gln	Asn	Met
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Ile	Val	Glu	Glu	Cys	Gly	Cys	Ser								
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 <211> 424
 <212> PRT

<213> Rattus norvegicus

<400> 8

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			20					25					30			
Pro	Asp	Cys	Pro	Ser	Cys	Ala	Leu	Ala	Thr	Leu	Pro	Lys	Asp	Gly	Pro	
		35						40					45			
Asn	Ser	Gln	Pro	Glu	Met	Val	Glu	Ala	Val	Lys	Lys	His	Ile	Leu	Asn	
	50						55					60				
Met	Leu	His	Leu	Lys	Lys	Arg	Pro	Asp	Val	Thr	Gln	Pro	Val	Pro	Lys	
65					70					75					80	
Ala	Ala	Leu	Leu	Asn	Ala	Ile	Arg	Lys	Leu	His	Val	Gly	Lys	Val	Gly	
				85					90						95	
Glu	Asn	Gly	Tyr	Val	Glu	Ile	Glu	Asp	Asp	Ile	Gly	Arg	Arg	Ala	Glu	
			100					105					110			
Met	Asn	Glu	Leu	Met	Glu	Gln	Thr	Ser	Glu	Ile	Ile	Thr	Phe	Ala	Glu	
		115						120					125			
Ser	Gly	Thr	Ala	Arg	Lys	Thr	Leu	His	Phe	Glu	Ile	Ser	Lys	Glu	Gly	
	130						135					140				
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145					150						155				160	
Val	Pro	Lys	Ala	Asn	Arg	Thr	Arg	Thr	Lys	Val	Thr	Ile	Arg	Leu	Phe	
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Gln	Gln	Gln	Lys	His	Pro	Gln	Gly	Ser	Leu	Asp	Met	Gly	Asp	Glu	Ala	
			180						185				190			
Glu	Glu	Met	Gly	Leu	Lys	Gly	Glu	Arg	Ser	Glu	Leu	Leu	Leu	Ser	Glu	
		195						200					205			
Lys	Val	Val	Asp	Ala	Arg	Lys	Ser	Thr	Trp	His	Ile	Phe	Pro	Val	Ser	
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Ser	Ser	Ile	Gln	Arg	Leu	Leu	Asp	Gln	Gly	Lys	Ser	Ser	Leu	Asp	Val	
225					230					235					240	
Arg	Ile	Ala	Cys	Glu	Gln	Cys	Gln	Glu	Ser	Gly	Ala	Ser	Leu	Val	Leu	
				245						250					255	
Leu	Gly	Lys	Lys	Lys	Lys	Lys	Glu	Val	Asp	Gly	Asp	Gly	Lys	Lys	Lys	
			260					265					270			
Asp	Gly	Ser	Asp	Gly	Gly	Leu	Glu	Glu	Glu	Lys	Glu	Gln	Ser	His	Arg	
			275					280					285			

Pro Phe Leu Met Leu Gln Ala Arg Gln Ser Glu Asp His Pro His Arg